

SEQUENCE LISTING

<110> INCYTE GENOMICS, INC.
 TANG, Y. Tom
 HILLMAN, Jennifer L.
 BANDMAN, Olga
 YUE, Henry
 BAUGHN, Mariah R.
 LAL, Preeti
 LU, Dyung Aina M.
 SHAH, Purvi
 AZIMZAI, Yalda

<120> HUMAN SYNTHETASES

<130> PF-0721 PCT

<140> To Be Assigned

<141> Herewith

<150> 60/144,992; 60,168,858

<151> 1999-07-22; 1999-12-02

<160> 30

<170> PERL Program

<210> 1

<211> 1176

<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 1806212CD1

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Ile	Glu	Lys	Glu	Ile	Gln	Gln	Lys	Trp	Asp	Thr	Glu	Arg	Val	Phe
				20					25					30
Glu	Val	Asn	Ala	Ser	Asn	Leu	Glu	Lys	Gln	Thr	Ser	Lys	Gly	Lys
				35					40					45
Tyr	Phe	Val	Thr	Phe	Pro	Tyr	Pro	Tyr	Met	Asn	Gly	Arg	Leu	His
				50					55					60
Leu	Gly	His	Thr	Phe	Ser	Leu	Ser	Lys	Cys	Glu	Phe	Ala	Val	Gly
				65					70					75
Tyr	Gln	Arg	Leu	Lys	Gly	Lys	Cys	Cys	Leu	Phe	Pro	Phe	Gly	Leu
				80					85					90
His	Cys	Thr	Gly	Met	Pro	Ile	Lys	Ala	Cys	Ala	Asp	Lys	Leu	Lys
				95					100					105
Arg	Glu	Ile	Glu	Leu	Tyr	Gly	Cys	Pro	Pro	Asp	Phe	Pro	Asp	Glu
				110					115					120
Glu	Glu	Glu	Glu	Glu	Glu	Thr	Ser	Val	Lys	Thr	Glu	Asp	Ile	Ile
				125					130					135
Ile	Lys	Asp	Lys	Ala	Lys	Gly	Lys	Lys	Ser	Lys	Ala	Ala	Ala	Lys
				140					145					150
Ala	Gly	Ser	Ser	Lys	Tyr	Gln	Trp	Gly	Ile	Met	Lys	Ser	Leu	Gly
				155					160					165
Leu	Ser	Asp	Glu	Glu	Ile	Val	Lys	Phe	Ser	Glu	Ala	Glu	His	Trp
				170					175					180
Leu	Asp	Tyr	Phe	Pro	Pro	Leu	Ala	Ile	Gln	Asp	Leu	Lys	Arg	Met
				185					190					195

Gly	Leu	Lys	Val	Asp	Trp	Arg	Arg	Ser	Phe	Ile	Thr	Thr	Asp	Val
				200					205					210
Asn	Pro	Tyr	Tyr	Asp	Ser	Phe	Val	Arg	Trp	Gln	Phe	Leu	Thr	Leu
				215					220					225
Arg	Glu	Arg	Asn	Lys	Ile	Lys	Phe	Gly	Lys	Arg	Tyr	Thr	Ile	Tyr
				230					235					240
Ser	Pro	Lys	Asp	Gly	Gln	Pro	Cys	Met	Asp	His	Asp	Arg	Gln	Thr
				245					250					255
Gly	Glu	Gly	Val	Gly	Pro	Gln	Glu	Tyr	Thr	Leu	Leu	Lys	Leu	Lys
				260					265					270
Val	Leu	Glu	Pro	Tyr	Pro	Ser	Lys	Leu	Ser	Gly	Leu	Lys	Gly	Lys
				275					280					285
Asn	Ile	Phe	Leu	Val	Ala	Ala	Thr	Leu	Arg	Pro	Glu	Thr	Met	Phe
				290					295					300
Gly	Gln	Thr	Asn	Cys	Trp	Val	Arg	Pro	Asp	Met	Lys	Tyr	Ile	Gly
				305					310					315
Phe	Glu	Thr	Val	Asn	Gly	Asp	Ile	Phe	Ile	Cys	Thr	Gln	Lys	Ala
				320					325					330
Ala	Arg	Asn	Met	Ser	Tyr	Gln	Gly	Phe	Thr	Lys	Asp	Asn	Gly	Val
				335					340					345
Val	Pro	Val	Val	Lys	Glu	Leu	Met	Gly	Glu	Glu	Ile	Leu	Gly	Ala
				350					355					360
Ser	Leu	Ser	Ala	Pro	Leu	Thr	Ser	Tyr	Lys	Val	Ile	Tyr	Val	Leu
				365					370					375
Pro	Met	Leu	Thr	Ile	Lys	Glu	Asp	Lys	Gly	Thr	Gly	Val	Val	Thr
				380					385					390
Ser	Val	Pro	Ser	Asp	Ser	Pro	Asp	Asp	Ile	Ala	Ala	Leu	Arg	Asp
				395					400					405
Leu	Lys	Lys	Lys	Gln	Ala	Leu	Arg	Ala	Lys	Tyr	Gly	Ile	Arg	Asp
				410					415					420
Asp	Met	Val	Leu	Pro	Phe	Glu	Pro	Val	Pro	Val	Ile	Glu	Ile	Pro
				425					430					435
Gly	Phe	Gly	Asn	Leu	Ser	Ala	Val	Thr	Ile	Cys	Asp	Glu	Leu	Lys
				440					445					450
Ile	Gln	Ser	Gln	Asn	Asp	Arg	Glu	Lys	Leu	Ala	Glu	Ala	Lys	Glu
				455					460					465
Lys	Ile	Tyr	Leu	Lys	Gly	Phe	Tyr	Glu	Gly	Ile	Met	Leu	Val	Asp
				470					475					480
Gly	Phe	Lys	Gly	Gln	Lys	Val	Gln	Asp	Val	Lys	Lys	Thr	Ile	Gln
				485					490					495
Lys	Lys	Met	Ile	Asp	Ala	Gly	Asp	Ala	Leu	Ile	Tyr	Met	Glu	Pro
				500					505					510
Glu	Lys	Gln	Val	Met	Ser	Arg	Ser	Ser	Asp	Glu	Cys	Val	Val	Ala
				515					520					525
Leu	Cys	Asp	Gln	Trp	Tyr	Leu	Asp	Tyr	Gly	Glu	Glu	Asn	Trp	Lys
				530					535					540
Lys	Gln	Thr	Ser	Gln	Cys	Leu	Lys	Asn	Leu	Glu	Thr	Phe	Cys	Glu
				545					550					555
Glu	Thr	Arg	Arg	Asn	Phe	Glu	Ala	Thr	Leu	Gly	Trp	Leu	Gln	Glu
				560					565					570
His	Ala	Cys	Ser	Arg	Thr	Tyr	Gly	Leu	Gly	Thr	His	Leu	Pro	Trp
				575					580					585
Asp	Glu	Gln	Trp	Leu	Ile	Glu	Ser	Leu	Ser	Asp	Ser	Thr	Ile	Tyr
				590					595					600
Met	Ala	Phe	Tyr	Thr	Val	Ala	His	Leu	Leu	Gln	Gly	Gly	Asn	Leu
				605					610					615
His	Gly	Gln	Ala	Glu	Ser	Pro	Leu	Gly	Ile	Arg	Pro	Gln	Gln	Met
				620					625					630
Thr	Lys	Glu	Val	Trp	Asp	Tyr	Val	Phe	Phe	Lys	Glu	Ala	Pro	Phe
				635					640					645
Pro	Lys	Thr	Gln	Ile	Ala	Lys	Glu	Lys	Leu	Asp	Gln	Leu	Lys	Gln
				650					655					660
Glu	Phe	Glu	Phe	Trp	Tyr	Pro	Val	Asp	Leu	Arg	Val	Ser	Gly	Lys
				665					670					675
Asp	Leu	Val	Pro	Asn	His	Leu	Ser	Tyr	Tyr	Leu	Tyr	Asn	His	Val

	680		685		690
Ala Met Trp Pro	Glu Gln Ser Asp Lys	Trp Pro Thr Ala Val	Arg		
	695		700		705
Ala Asn Gly His	Leu Leu Leu Asn Ser	Glu Lys Met Ser Lys	Ser		
	710		715		720
Thr Gly Asn Phe	Leu Thr Leu Thr Gln	Ala Ile Asp Lys Phe	Ser		
	725		730		735
Ala Asp Gly Met	Arg Leu Ala Leu Ala	Asp Ala Gly Asp Thr	Val		
	740		745		750
Glu Asp Ala Asn	Phe Val Glu Ala Met	Ala Asp Ala Gly Ile	Leu		
	755		760		765
Arg Leu Tyr Thr	Trp Val Glu Trp Val	Lys Glu Met Val Ala	Asn		
	770		775		780
Trp Asp Ser Leu	Arg Ser Gly Pro Ala	Ser Thr Phe Asn Asp	Arg		
	785		790		795
Val Phe Ala Ser	Glu Leu Asn Ala Gly	Ile Ile Lys Thr Asp	Gln		
	800		805		810
Asn Tyr Glu Lys	Met Met Phe Lys Glu	Ala Leu Lys Thr Gly	Phe		
	815		820		825
Phe Glu Phe Gln	Ala Ala Lys Asp Lys	Trp Arg Glu Leu Ala	Val		
	830		835		840
Glu Gly Met His	Arg Glu Leu Val Phe	Arg Phe Ile Glu Val	Gln		
	845		850		855
Thr Leu Leu Leu	Ala Pro Phe Cys Pro	His Leu Cys Glu His	Ile		
	860		865		870
Trp Thr Leu Leu	Gly Lys Pro Asp Ser	Ile Met Asn Ala Ser	Trp		
	875		880		885
Pro Val Ala Gly	Pro Val Asn Glu Val	Leu Ile His Ser Ser	Gln		
	890		895		900
Tyr Leu Met Glu	Val Thr His Asp Leu	Arg Leu Arg Leu Lys	Asn		
	905		910		915
Tyr Met Met Pro	Ala Lys Gly Lys Lys	Thr Asp Lys Gln Pro	Leu		
	920		925		930
Gln Lys Pro Ser	His Cys Thr Ile Tyr	Val Ala Lys Asn Tyr	Pro		
	935		940		945
Pro Trp Gln His	Thr Thr Leu Ser Val	Leu Arg Lys His Phe	Glu		
	950		955		960
Ala Asn Asn Gly	Lys Leu Pro Asp Asn	Lys Val Ile Ala Ser	Glu		
	965		970		975
Leu Gly Ser Met	Pro Glu Leu Lys Lys	Tyr Met Lys Lys Val	Met		
	980		985		990
Pro Phe Val Ala	Met Ile Lys Glu Asn	Leu Glu Lys Met Gly	Pro		
	995		1000		1005
Arg Ile Leu Asp	Leu Gln Leu Glu Phe	Asp Glu Lys Ala Val	Leu		
	1010		1015		1020
Met Glu Asn Ile	Val Tyr Leu Thr Asn	Ser Leu Glu Leu Glu	His		
	1025		1030		1035
Ile Glu Val Lys	Phe Ala Ser Glu Ala	Glu Asp Lys Ile Arg	Glu		
	1040		1045		1050
Asp Cys Cys Pro	Gly Lys Pro Leu Asn	Val Phe Arg Ile Glu	Pro		
	1055		1060		1065
Gly Val Ser Val	Ser Leu Val Asn Pro	Gln Pro Ser Asn Gly	His		
	1070		1075		1080
Phe Ser Thr Lys	Ile Glu Ile Arg Gln	Gly Asp Asn Cys Asp	Ser		
	1085		1090		1095
Ile Ile Arg Arg	Leu Met Lys Met Asn	Arg Gly Ile Lys Asp	Leu		
	1100		1105		1110
Ser Lys Val Lys	Leu Met Arg Phe Asp	Asp Pro Leu Leu Gly	Pro		
	1115		1120		1125
Arg Arg Val Pro	Val Leu Gly Lys Glu	Tyr Thr Glu Lys Thr	Pro		
	1130		1135		1140
Ile Ser Glu His	Ala Val Phe Asn Val	Asp Leu Met Ser Lys	Lys		
	1145		1150		1155
Ile His Leu Thr	Glu Asn Gly Ile Arg	Val Asp Ile Gly Asp	Thr		
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Ile Ile Tyr Leu Val His
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<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 2083883CD1

<400> 2

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Gly	Lys	Lys	Asp	Arg	Asp	Ser	Cys	Gly	Arg	Lys	Asn	Ser	Glu	Pro	20	25	30	35
Gly	Ser	Pro	His	Ser	Leu	Glu	Ala	Leu	Arg	Asp	Ala	Ala	Pro	Ser	40	45	50	55
Gln	Gly	Leu	Asn	Phe	Leu	Leu	Leu	Phe	Thr	Lys	Met	Leu	Phe	Ile	60	65	70	75
Phe	Asn	Phe	Leu	Phe	Ser	Pro	Leu	Pro	Thr	Pro	Ala	Leu	Ile	Cys	80	85	90	95
Ile	Leu	Thr	Phe	Gly	Ala	Ala	Ile	Phe	Leu	Trp	Leu	Ile	Thr	Arg	100	105	110	115
Pro	Gln	Pro	Val	Leu	Pro	Leu	Leu	Asp	Leu	Asn	Asn	Gln	Ser	Val	120	125	130	135
Gly	Ile	Glu	Gly	Gly	Ala	Arg	Lys	Gly	Val	Ser	Gln	Lys	Asn	Asn	140	145	150	155
Asp	Leu	Thr	Ser	Cys	Cys	Phe	Ser	Asp	Ala	Lys	Thr	Met	Tyr	Glu	160	165	170	175
Val	Phe	Gln	Arg	Gly	Leu	Ala	Val	Ser	Asp	Asn	Gly	Pro	Cys	Leu	180	185	190	195
Gly	Tyr	Arg	Lys	Pro	Asn	Gln	Pro	Tyr	Arg	Trp	Leu	Ser	Tyr	Lys	200	205	210	215
Gln	Val	Ser	Asp	Arg	Ala	Glu	Tyr	Leu	Gly	Ser	Cys	Leu	Leu	His	220	225	230	235
Lys	Gly	Tyr	Lys	Ser	Ser	Pro	Asp	Gln	Phe	Val	Gly	Ile	Phe	Ala	240	245	250	255
Gln	Asn	Arg	Pro	Glu	Trp	Ile	Ile	Ser	Glu	Leu	Ala	Cys	Tyr	Thr	260	265	270	275
Tyr	Ser	Met	Val	Ala	Val	Pro	Leu	Tyr	Asp	Thr	Leu	Gly	Pro	Glu	280	285	290	295
Ala	Ile	Val	His	Ile	Val	Asn	Lys	Ala	Asp	Ile	Ala	Val	Val	Ile	300	305	310	315
Cys	Asp	Thr	Pro	Gln	Lys	Ala	Leu	Val	Leu	Ile	Gly	Asn	Val	Glu	320	325	330	335
Lys	Gly	Phe	Thr	Pro	Ser	Leu	Lys	Val	Ile	Ile	Leu	Met	Asp	Pro	340	345	350	355
Phe	Asp	Asp	Asp	Leu	Lys	Gln	Arg	Gly	Glu	Lys	Ser	Gly	Ile	Glu	360	365	370	375
Ile	Leu	Ser	Leu	Tyr	Asp	Ala	Glu	Asn	Leu	Gly	Lys	Glu	His	Phe	380	385	390	395
Arg	Lys	Pro	Val	Pro	Pro	Ser	Pro	Glu	Asp	Leu	Ser	Val	Ile	Cys	400	405	410	415
Phe	Thr	Ser	Gly	Thr	Thr	Gly	Asp	Pro	Lys	Gly	Ala	Met	Ile	Thr	420	425	430	435
His	Gln	Asn	Ile	Val	Ser	Asn	Ala	Ala	Ala	Phe	Leu	Lys	Cys	Val	440	445	450	455
Glu	His	Ala	Tyr	Glu	Pro	Thr	Pro	Asp	Asp	Val	Ala	Ile	Ser	Tyr	460	465	470	475
Leu	Pro	Leu	Ala	His	Met	Phe	Glu	Arg	Ile	Val	Gln	Ala	Val	Val	480	485	490	495
Tyr	Ser	Cys	Gly	Ala	Arg	Val	Gly	Phe	Phe	Gln	Gly	Asp	Ile	Arg	500	505	510	515
Leu	Leu	Ala	Asp	Asp	Met	Lys	Thr	Leu	Lys	Pro	Thr	Leu	Phe	Pro	520	525	530	535

Ala Val Pro Arg	395	Leu Leu Asn Arg Ile	400	Tyr Asp Lys Val Gln Asn	405
	410		415		420
Glu Ala Lys Thr	425	Leu Lys Lys Phe	430	Leu Leu Lys Leu Ala Val	435
Ser Ser Lys Phe	440	Lys Glu Leu Gln Lys	445	Gly Ile Ile Arg His Asp	450
Ser Phe Trp Asp	455	Lys Leu Ile Phe Ala	460	Lys Ile Gln Asp Ser Leu	465
Gly Gly Arg Val	470	Arg Val Ile Val Thr	475	Gly Ala Ala Pro Met Ser	480
Thr Ser Val Met	485	Thr Phe Phe Arg Ala	490	Ala Met Gly Cys Gln Val	495
Tyr Glu Ala Tyr	500	Gly Gln Thr Glu Cys	505	Thr Gly Gly Cys Thr Phe	510
Thr Leu Pro Gly	515	Asp Trp Thr Ser Gly	520	His Val Gly Val Pro Leu	525
Ala Cys Asn Tyr	530	Val Lys Leu Glu Asp	535	Val Ala Asp Met Asn Tyr	540
Phe Thr Val Asn	545	Asn Glu Gly Glu Val	550	Cys Ile Lys Gly Thr Asn	555
Val Phe Lys Gly	560	Tyr Leu Lys Asp Pro	565	Glu Lys Thr Gln Glu Ala	570
Leu Asp Ser Asp	575	Gly Trp Leu His Thr	580	Gly Asp Ile Gly Arg Trp	585
Leu Pro Asn Gly	590	Thr Leu Lys Ile Ile	595	Asp Arg Lys Lys Asn Ile	600
Phe Lys Leu Ala	605	Gln Gly Glu Tyr Ile	610	Ala Pro Glu Lys Ile Glu	615
Asn Ile Tyr Asn	620	Arg Ser Gln Pro Val	625	Leu Gln Ile Phe Val His	630
Gly Glu Ser Leu	635	Arg Ser Ser Leu Val	640	Gly Val Val Val Pro Asp	645
Thr Asp Val Leu	650	Pro Ser Phe Ala Ala	655	Lys Leu Gly Val Lys Gly	660
Ser Phe Glu Glu	665	Leu Cys Gln Asn Gln	670	Val Val Arg Glu Ala Ile	675
Leu Glu Asp Leu	680	Gln Lys Ile Gly Lys	685	Glu Ser Gly Leu Lys Thr	690
Phe Glu Gln Val	695	Lys Ala Ile Phe Leu	700	His Pro Glu Pro Phe Ser	705
Ile Glu Asn Gly	710	Leu Leu Thr Pro Thr	715	Leu Lys Ala Lys Arg Gly	720
Glu Leu Ser Lys	725	Tyr Phe Arg Thr Gln	730	Ile Asp Ser Leu Tyr Glu	735
His Ile Gln Asp					

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<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 2454288CD1

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Gly Arg Thr Tyr	20	Thr Asp Glu Glu Phe	25	Asp Glu Leu Cys Phe Glu	30
Phe Gly Leu Glu	35	Leu Asp Glu Ile Thr	40	Ser Glu Lys Glu Ile Ile	45
Ser Lys Glu Gln	50	Gly Asn Val Lys Ala	55	Ala Gly Ala Ser Asp Val	60

Val	Leu	Tyr	Lys	Ile	Asp	Val	Pro	Ala	Asn	Arg	Tyr	Asp	Leu	Leu
				65					70					75
Cys	Leu	Glu	Gly	Leu	Val	Arg	Gly	Leu	Gln	Val	Phe	Lys	Glu	Arg
				80					85					90
Ile	Lys	Ala	Pro	Val	Tyr	Lys	Arg	Val	Met	Pro	Asp	Gly	Lys	Ile
				95					100					105
Gln	Lys	Leu	Ile	Ile	Thr	Glu	Glu	Thr	Ala	Lys	Ile	Arg	Pro	Phe
				110					115					120
Ala	Val	Ala	Ala	Val	Leu	Arg	Asn	Ile	Lys	Phe	Thr	Lys	Asp	Arg
				125					130					135
Tyr	Asp	Ser	Phe	Ile	Glu	Leu	Gln	Glu	Lys	Leu	His	Gln	Asn	Ile
				140					145					150
Cys	Arg	Lys	Arg	Ala	Leu	Val	Ala	Ile	Gly	Thr	His	Asp	Leu	Asp
				155					160					165
Thr	Leu	Ser	Gly	Pro	Phe	Thr	Tyr	Thr	Ala	Lys	Arg	Pro	Ser	Asp
				170					175					180
Ile	Lys	Phe	Lys	Pro	Leu	Asn	Lys	Thr	Lys	Glu	Tyr	Thr	Ala	Cys
				185					190					195
Glu	Leu	Met	Asn	Ile	Tyr	Lys	Thr	Asp	Asn	His	Leu	Lys	His	Tyr
				200					205					210
Leu	His	Ile	Ile	Glu	Asn	Lys	Pro	Leu	Tyr	Pro	Val	Ile	Tyr	Asp
				215					220					225
Ser	Asn	Gly	Val	Val	Leu	Ser	Met	Pro	Pro	Ile	Ile	Asn	Gly	Asp
				230					235					240
His	Ser	Arg	Ile	Thr	Val	Asn	Thr	Arg	Asn	Ile	Phe	Ile	Glu	Cys
				245					250					255
Thr	Gly	Thr	Asp	Phe	Thr	Lys	Ala	Lys	Ile	Val	Leu	Asp	Ile	Ile
				260					265					270
Val	Thr	Met	Phe	Ser	Glu	Tyr	Cys	Glu	Asn	Gln	Phe	Thr	Val	Glu
				275					280					285
Ala	Ala	Glu	Val	Val	Phe	Pro	Asn	Gly	Lys	Ser	His	Thr	Phe	Pro
				290					295					300
Glu	Leu	Ala	Tyr	Arg	Lys	Glu	Met	Val	Arg	Ala	Asp	Leu	Ile	Asn
				305					310					315
Lys	Lys	Val	Gly	Ile	Arg	Glu	Thr	Pro	Glu	Asn	Leu	Ala	Lys	Leu
				320					325					330
Leu	Thr	Arg	Met	Tyr	Leu	Lys	Ser	Glu	Val	Ile	Gly	Asp	Gly	Asn
				335					340					345
Gln	Ile	Glu	Ile	Glu	Ile	Pro	Pro	Thr	Arg	Ala	Asp	Ile	Ile	His
				350					355					360
Ala	Cys	Asp	Ile	Val	Glu	Asp	Ala	Ala	Ile	Ala	Tyr	Gly	Tyr	Asn
				365					370					375
Asn	Ile	Gln	Met	Thr	Leu	Pro	Lys	Thr	Tyr	Thr	Ile	Ala	Asn	Gln
				380					385					390
Phe	Pro	Leu	Asn	Lys	Leu	Thr	Glu	Leu	Leu	Arg	His	Asp	Met	Ala
				395					400					405
Ala	Ala	Gly	Phe	Thr	Glu	Ala	Leu	Thr	Phe	Ala	Leu	Cys	Ser	Gln
				410					415					420
Glu	Asp	Ile	Ala	Asp	Lys	Leu	Gly	Val	Asp	Ile	Ser	Ala	Thr	Lys
				425					430					435
Ala	Val	His	Ile	Ser	Asn	Pro	Lys	Thr	Ala	Glu	Phe	Gln	Val	Ala
				440					445					450
Arg	Thr	Thr	Leu	Leu	Pro	Gly	Leu	Leu	Lys	Thr	Ile	Ala	Ala	Asn
				455					460					465
Arg	Lys	Met	Pro	Leu	Pro	Leu	Lys	Leu	Phe	Glu	Ile	Ser	Asp	Ile
				470					475					480
Val	Ile	Lys	Asp	Ser	Asn	Thr	Asp	Val	Gly	Ala	Lys	Asn	Tyr	Arg
				485					490					495
His	Leu	Cys	Ala	Val	Tyr	Tyr	Asn	Lys	Asn	Pro	Gly	Phe	Glu	Ile
				500					505					510
Ile	His	Gly	Leu	Leu	Asp	Arg	Ile	Met	Gln	Leu	Leu	Asp	Val	Pro
				515					520					525
Pro	Gly	Glu	Asp	Lys	Gly	Gly	Tyr	Val	Ile	Lys	Ala	Ser	Glu	Gly
				530					535					540
Pro	Ala	Phe	Phe	Pro	Gly	Arg	Cys	Ala	Glu	Ile	Phe	Ala	Arg	Gly

Gln Ser Val Gly	545	Leu Gly Val Leu	550	Pro Asp Val Ile	555
Lys Phe Glu Leu	560	Thr Met Pro Cys Ser	565	Ser Leu Glu Ile Asn	570
Gly Pro Phe Leu	575		580		585

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<220>
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 <223> Incyte ID No: 1513539CD1

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Cys Tyr Asn Val Thr	Asn Ile Gly Phe	Lys Ser Pro Ser Asp	Phe
20	25		30
Trp Gln Ser Val His	Ser Thr Leu Pro Arg	Glu Leu Ala Pro	Cys
35	40		45
Leu Val Phe Asn Thr	Ser Pro Asn Leu	Ala Leu Phe Ser	Ala
50	55		60
Phe Ala Phe Ile Val	Val Lys Asp Ser	Ala Gly Asp Ser	Val
65	70		75
Val Val Gln Glu Leu	Lys Ser Met Val	Ala Thr Lys Ile	Lys
80	85		90
Tyr Ala Val Pro Asp	Glu Ile Leu Val	Val Lys Arg Leu	Pro
95	100		105
Thr Arg Ser Gly Lys	Val Met Arg Arg	Leu Leu Arg Lys	Ile
110	115		120
Thr Ser Glu Ala Gln	Glu Leu Gly Asp	Thr Thr Leu Glu	Asp
125	130		135
Pro Ser Ile Ile Ala	Glu Ile Leu Ser	Val Tyr Gln Lys	Cys
140	145		150
Asp Lys Gln Ala Ala	Lys		
155			

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<220>
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 <223> Incyte ID No: 2148623CD1

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Ser Gln Asp Leu Lys	Glu Asp Leu Leu	Tyr Asn Leu Lys	Gln
20	25		30
Gly Pro Asn Ser Ser	Lys Gln Leu Leu	Lys Ser Asp Val	Asn
35	40		45
Gln Cys Leu Phe Ser	Ala His Val Leu	His Leu Arg Gly	Val
50	55		60
Thr Thr Gln Pro Val	Glu Asp Glu Arg	Gly Asn Val Phe	Leu
65	70		75
Asn Gly Glu Ile Phe	Ser Gly Ile Lys	Val Glu Ala Glu	Glu
80	85		90
Asp Thr Gln Ile Leu	Phe Asn Tyr Leu	Ser Ser Cys Lys	Asn
95	100		105
Ser Glu Ile Leu Ser	Leu Phe Ser Glu	Val Gln Gly Pro	Trp
110	115		120

Phe	Ile	Tyr	Tyr	Gln	Ala	Ser	Ser	His	Tyr	Leu	Trp	Phe	Gly	Arg
				125					130					135
Asp	Phe	Phe	Gly	Arg	Arg	Ser	Leu	Leu	Trp	His	Phe	Ser	Asn	Leu
				140					145					150
Gly	Lys	Ser	Phe	Cys	Leu	Ser	Ser	Val	Gly	Thr	Gln	Thr	Ser	Gly
				155					160					165
Leu	Ala	Asn	Gln	Trp	Gln	Glu	Val	Pro	Ala	Ser	Gly	Leu	Phe	Arg
				170					175					180
Ile	Asp	Leu	Lys	Ser	Thr	Val	Ile	Ser	Arg	Cys	Ile	Ile	Leu	Gln
				185					190					195
Leu	Tyr	Pro	Trp	Lys	Tyr	Ile	Ser	Arg	Glu	Asn	Ile	Ile	Glu	Glu
				200					205					210
Asn	Val	Asn	Ser	Leu	Ser	Gln	Ile	Ser	Ala	Asp	Leu	Pro	Ala	Phe
				215					220					225
Val	Ser	Val	Val	Ala	Asn	Glu	Ala	Lys	Leu	Tyr	Leu	Glu	Lys	Pro
				230					235					240
Val	Val	Pro	Leu	Asn	Met	Met	Leu	Pro	Gln	Ala	Ala	Leu	Glu	Thr
				245					250					255
His	Cys	Ser	Asn	Ile	Ser	Asn	Val	Pro	Pro	Thr	Arg	Glu	Ile	Leu
				260					265					270
Gln	Val	Phe	Leu	Thr	Asp	Val	His	Met	Lys	Glu	Val	Ile	Gln	Gln
				275					280					285
Phe	Ile	Asp	Val	Leu	Ser	Val	Ala	Val	Lys	Lys	Arg	Val	Leu	Cys
				290					295					300
Leu	Pro	Arg	Asp	Glu	Asn	Leu	Thr	Ala	Asn	Glu	Val	Leu	Lys	Thr
				305					310					315
Cys	Asp	Arg	Lys	Ala	Asn	Val	Ala	Ile	Leu	Phe	Ser	Gly	Gly	Ile
				320					325					330
Asp	Ser	Met	Val	Ile	Ala	Thr	Leu	Ala	Asp	Arg	His	Ile	Pro	Leu
				335					340					345
Asp	Glu	Pro	Ile	Asp	Leu	Leu	Asn	Val	Ala	Phe	Ile	Ala	Glu	Glu
				350					355					360
Lys	Thr	Met	Pro	Thr	Thr	Phe	Asn	Arg	Glu	Gly	Asn	Lys	Gln	Lys
				365					370					375
Asn	Lys	Cys	Glu	Ile	Pro	Ser	Glu	Glu	Phe	Ser	Lys	Asp	Val	Ala
				380					385					390
Ala	Ala	Ala	Ala	Asp	Ser	Pro	Asn	Lys	His	Val	Ser	Val	Pro	Asp
				395					400					405
Arg	Ile	Thr	Gly	Arg	Ala	Gly	Leu	Lys	Glu	Leu	Gln	Ala	Val	Ser
				410					415					420
Pro	Ser	Arg	Ile	Trp	Asn	Phe	Val	Glu	Ile	Asn	Val	Ser	Met	Glu
				425					430					435
Glu	Leu	Gln	Lys	Leu	Arg	Arg	Thr	Arg	Ile	Cys	His	Leu	Ile	Arg
				440					445					450
Pro	Leu	Asp	Thr	Val	Leu	Asp	Asp	Ser	Ile	Gly	Cys	Ala	Val	Trp
				455					460					465
Phe	Ala	Ser	Arg	Gly	Ile	Gly	Trp	Leu	Val	Ala	Gln	Glu	Gly	Val
				470					475					480
Lys	Ser	Tyr	Gln	Ser	Asn	Ala	Lys	Val	Val	Leu	Thr	Gly	Ile	Gly
				485					490					495
Ala	Asp	Glu	Gln	Leu	Ala	Gly	Tyr	Ser	Arg	His	Arg	Val	Arg	Phe
				500					505					510
Gln	Ser	His	Gly	Leu	Glu	Gly	Leu	Asn	Lys	Glu	Ile	Met	Met	Glu
				515					520					525
Leu	Gly	Arg	Ile	Ser	Ser	Arg	Asn	Leu	Gly	Arg	Asp	Asp	Arg	Val
				530					535					540
Ile	Gly	Asp	His	Gly	Lys	Glu	Ala	Arg	Phe	Pro	Phe	Leu	Asp	Glu
				545					550					555
Asn	Val	Val	Ser	Phe	Leu	Asn	Ser	Leu	Pro	Ile	Trp	Glu	Lys	Ala
				560					565					570
Asn	Leu	Thr	Leu	Pro	Arg	Gly	Ile	Gly	Glu	Lys	Leu	Leu	Leu	Arg
				575					580					585
Leu	Ala	Ala	Val	Glu	Leu	Gly	Leu	Thr	Ala	Ser	Ala	Leu	Leu	Pro
				590					595					600
Lys	Arg	Ala	Met	Gln	Phe	Gly	Ser	Arg	Ile	Ala	Lys	Met	Glu	Lys

Ile Asn Glu Lys	605	Ser Asp Lys Cys	610	Gly Arg Leu Gln Ile	615
	620		625		630
Ser Leu Glu Asn Leu	635	Ser Ile Glu Lys	640	Glu Thr Lys Leu	

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Glu Glu Glu Ile Lys	20	Ser Leu Thr Ala	25	Glu Ile Asp Arg Leu Lys	30
Asn Cys Gly Cys Leu	35	Gly Ala Ser Pro	40	Asn Leu Glu Gln Leu Gln	45
Glu Glu Asn Leu Lys	50	Leu Lys Tyr Arg	55	Leu Asn Ile Leu Arg Lys	60
Ser Leu Gln Ala Glu	65	Arg Asn Lys Pro	70	Thr Lys Asn Met Ile Asn	75
Ile Ile Ser Arg Leu	80	Gln Glu Val Phe	85	Gly His Ala Ile Lys Ala	90
Ala Tyr Pro Asp Leu	95	Glu Asn Pro Pro	100	Leu Leu Val Thr Pro Ser	105
Gln Gln Ala Lys Phe	110	Gly Asp Tyr Gln	115	Cys Asn Ser Ala Met Gly	120
Ile Ser Gln Met Leu	125	Lys Thr Lys Glu	130	Gln Lys Val Asn Pro Arg	135
Glu Ile Ala Glu Asn	140	Ile Thr Lys His	145	Leu Pro Asp Asn Glu Cys	150
Ile Glu Lys Val Glu	155	Ile Ala Gly Pro	160	Gly Phe Ile Asn Val His	165
Leu Arg Lys Asp Phe	170	Val Ser Glu Gln	175	Leu Thr Ser Leu Leu Val	180
Asn Gly Val Gln Leu	185	Pro Ala Leu Gly	190	Glu Asn Lys Lys Val Ile	195
Val Asp Phe Ser Ser	200	Pro Asn Ile Ala	205	Lys Glu Met His Val Gly	210
His Leu Arg Ser Thr	215	Ile Ile Gly Glu	220	Ser Ile Ser Arg Leu Phe	225
Glu Phe Ala Gly Tyr	230	Asp Val Leu Arg	235	Leu Asn His Val Gly Asp	240
Trp Gly Thr Gln Phe	245	Gly Met Leu Ile	250	Ala His Leu Gln Asp Lys	255
Phe Pro Asp Tyr Leu	260	Thr Val Ser Pro	265	Pro Ile Gly Asp Leu Gln	270
Val Phe Tyr Lys Glu	275	Ser Lys Lys Arg	280	Phe Asp Thr Glu Glu Glu	285
Phe Lys Lys Arg Ala	290	Tyr Gln Cys Val	295	Val Leu Leu Gln Gly Lys	300
Asn Pro Asp Ile Thr	305	Lys Ala Trp Lys	310	Leu Ile Cys Asp Val Ser	315
Arg Gln Glu Leu Asn	320	Lys Ile Tyr Asp	325	Ala Leu Asp Val Ser Leu	330
Ile Glu Arg Gly Glu	335	Ser Phe Tyr Gln	340	Asp Arg Met Asn Asp Ile	345
Val Lys Glu Phe Glu	350	Asp Arg Gly Phe	355	Val Gln Val Asp Asp Gly	360
Arg Lys Ile Val Phe	365	Val Pro Gly Cys	370	Ser Ile Pro Leu Thr Ile	375

Val	Lys	Ser	Asp	Gly	Gly	Tyr	Thr	Tyr	Asp	Thr	Ser	Asp	Leu	Ala
				380					385					390
Ala	Ile	Lys	Gln	Arg	Leu	Phe	Glu	Glu	Lys	Ala	Asp	Met	Ile	Ile
				395					400					405
Tyr	Val	Val	Asp	Asn	Gly	Gln	Ser	Val	His	Phe	Gln	Thr	Ile	Phe
				410					415					420
Ala	Ala	Ala	Gln	Met	Ile	Gly	Trp	Tyr	Asp	Pro	Lys	Val	Thr	Arg
				425					430					435
Val	Phe	His	Ala	Gly	Phe	Gly	Val	Val	Leu	Gly	Glu	Asp	Lys	Lys
				440					445					450
Lys	Phe	Lys	Thr	Arg	Ser	Gly	Glu	Thr	Val	Arg	Leu	Met	Asp	Leu
				455					460					465
Leu	Gly	Glu	Gly	Leu	Lys	Arg	Ser	Met	Asp	Lys	Leu	Lys	Glu	Lys
				470					475					480
Glu	Arg	Asp	Lys	Val	Leu	Thr	Ala	Glu	Glu	Leu	Asn	Ala	Ala	Gln
				485					490					495
Thr	Ser	Val	Ala	Tyr	Gly	Cys	Ile	Lys	Tyr	Ala	Asp	Leu	Ser	His
				500					505					510
Asn	Arg	Leu	Asn	Asp	Tyr	Ile	Phe	Ser	Phe	Asp	Lys	Met	Leu	Asp
				515					520					525
Asp	Arg	Gly	Asn	Thr	Ala	Ala	Tyr	Leu	Leu	Tyr	Ala	Phe	Thr	Arg
				530					535					540
Ile	Arg	Ser	Ile	Ala	Arg	Leu	Ala	Asn	Ile	Asp	Glu	Glu	Met	Leu
				545					550					555
Gln	Lys	Ala	Ala	Arg	Glu	Thr	Lys	Ile	Leu	Leu	Asp	His	Glu	Lys
				560					565					570
Glu	Trp	Lys	Leu	Gly	Arg	Cys	Ile	Leu	Arg	Phe	Pro	Glu	Ile	Leu
				575					580					585
Gln	Lys	Ile	Leu	Asp	Asp	Leu	Phe	Leu	His	Thr	Leu	Cys	Asp	Tyr
				590					595					600
Ile	Tyr	Glu	Leu	Ala	Thr	Ala	Phe	Thr	Glu	Phe	Tyr	Asp	Ser	Cys
				605					610					615
Tyr	Cys	Val	Glu	Lys	Asp	Arg	Gln	Thr	Gly	Lys	Ile	Leu	Lys	Val
				620					625					630
Asn	Met	Trp	Arg	Met	Leu	Leu	Cys	Glu	Ala	Val	Ala	Ala	Val	Met
				635					640					645
Ala	Lys	Gly	Phe	Asp	Ile	Leu	Gly	Ile	Lys	Pro	Val	Gln	Arg	Met
				650					655					660

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<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 2662427CD1

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Met	Ala	Ala	Ala	Ser	Ala	Val	Ser	Val	Leu	Leu	Val	Ala	Ala	Glu
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Arg	Asn	Arg	Trp	His	Arg	Leu	Pro	Ser	Leu	Leu	Leu	Pro	Pro	Arg
				20					25					30
Thr	Trp	Val	Trp	Arg	Gln	Arg	Thr	Met	Lys	Tyr	Thr	Thr	Ala	Thr
				35					40					45
Gly	Arg	Asn	Ile	Thr	Lys	Val	Leu	Ile	Ala	Asn	Arg	Gly	Glu	Ile
				50					55					60
Ala	Cys	Arg	Val	Met	Arg	Thr	Ala	Lys	Lys	Leu	Gly	Val	Gln	Thr
				65					70					75
Val	Ala	Val	Tyr	Ser	Glu	Ala	Asp	Arg	Asn	Ser	Met	His	Val	Asp
				80					85					90
Met	Ala	Asp	Glu	Ala	Tyr	Ser	Ile	Gly	Pro	Ala	Pro	Ser	Gln	Gln
				95					100					105
Ser	Tyr	Leu	Ser	Met	Glu	Lys	Ile	Ile	Gln	Val	Ala	Lys	Thr	Ser
				110					115					120

Ala	Ala	Gln	Ala	Ile	His	Pro	Gly	Cys	Arg	Phe	Leu	Ser	Glu	Asn
				125					130					135
Met	Glu	Phe	Ala	Glu	Leu	Cys	Lys	Gln	Glu	Gly	Ile	Ile	Phe	Ile
				140					145					150
Gly	Pro	Pro	Pro	Ser	Ala	Ile	Arg	Asp	Met	Gly	Ile	Lys	Ser	Thr
				155					160					165
Ser	Lys	Ser	Ile	Met	Ala	Ala	Ala	Gly	Val	Pro	Val	Val	Glu	Gly
				170					175					180
Tyr	His	Gly	Glu	Asp	Gln	Ser	Asp	Gln	Cys	Leu	Lys	Glu	His	Ala
				185					190					195
Arg	Arg	Ile	Gly	Tyr	Pro	Val	Met	Ile	Lys	Ala	Val	Arg	Gly	Gly
				200					205					210
Gly	Gly	Lys	Gly	Met	Arg	Ile	Val	Arg	Ser	Glu	Gln	Glu	Phe	Gln
				215					220					225
Glu	Gln	Leu	Glu	Ser	Ala	Arg	Arg	Glu	Ala	Lys	Lys	Ser	Phe	Asn
				230					235					240
Asp	Asp	Ala	Met	Leu	Ile	Glu	Lys	Phe	Val	Asp	Thr	Pro	Arg	His
				245					250					255
Val	Glu	Val	Gln	Val	Phe	Gly	Asp	His	His	Gly	Asn	Ala	Val	Tyr
				260					265					270
Leu	Phe	Glu	Arg	Asp	Cys	Ser	Val	Gln	Arg	Arg	His	Gln	Lys	Ile
				275					280					285
Ile	Glu	Glu	Ala	Pro	Ala	Pro	Gly	Ile	Lys	Ser	Glu	Val	Arg	Lys
				290					295					300
Lys	Leu	Gly	Glu	Ala	Ala	Val	Arg	Ala	Ala	Lys	Ala	Val	Asn	Tyr
				305					310					315
Val	Gly	Ala	Gly	Thr	Val	Glu	Phe	Ile	Met	Asp	Ser	Lys	His	Asn
				320					325					330
Phe	Cys	Phe	Met	Glu	Met	Asn	Thr	Arg	Leu	Gln	Val	Glu	His	Pro
				335					340					345
Val	Thr	Glu	Met	Ile	Thr	Gly	Thr	Asp	Leu	Val	Glu	Trp	Gln	Leu
				350					355					360
Arg	Ile	Ala	Ala	Gly	Glu	Lys	Ile	Pro	Leu	Ser	Gln	Glu	Glu	Ile
				365					370					375
Thr	Leu	Gln	Gly	His	Ala	Phe	Glu	Ala	Arg	Ile	Tyr	Ala	Glu	Asp
				380					385					390
Pro	Ser	Asn	Asn	Phe	Met	Pro	Val	Ala	Gly	Pro	Leu	Val	His	Leu
				395					400					405
Ser	Thr	Pro	Arg	Ala	Asp	Pro	Ser	Thr	Arg	Ile	Glu	Thr	Gly	Val
				410					415					420
Arg	Gln	Gly	Asp	Glu	Val	Ser	Val	His	Tyr	Asp	Pro	Met	Ile	Ala
				425					430					435
Lys	Leu	Val	Val	Trp	Ala	Ala	Asp	Arg	Gln	Ala	Ala	Leu	Thr	Lys
				440					445					450
Leu	Arg	Tyr	Ser	Leu	Arg	Gln	Tyr	Asn	Ile	Val	Gly	Leu	Pro	Thr
				455					460					465
Asn	Ile	Asp	Phe	Leu	Leu	Asn	Leu	Ser	Gly	His	Pro	Glu	Phe	Glu
				470					475					480
Ala	Gly	Asn	Val	His	Thr	Asp	Phe	Ile	Pro	Gln	His	His	Lys	Gln
				485					490					495
Leu	Leu	Leu	Ser	Arg	Lys	Ala	Ala	Ala	Lys	Glu	Ser	Leu	Cys	Gln
				500					505					510
Ala	Ala	Leu	Gly	Leu	Ile	Leu	Lys	Glu	Lys	Ala	Met	Thr	Asp	Thr
				515					520					525
Phe	Thr	Leu	Gln	Ala	His	Asp	Gln	Phe	Ser	Pro	Phe	Ser	Ser	Ser
				530					535					540
Ser	Gly	Arg	Arg	Leu	Asn	Ile	Ser	Tyr	Thr	Arg	Asn	Met	Thr	Leu
				545					550					555
Lys	Asp	Gly	Lys	Asn	Asn	Val	Ala	Ile	Ala	Val	Thr	Tyr	Asn	His
				560					565					570
Asp	Gly	Ser	Tyr	Ser	Met	Gln	Ile	Glu	Asp	Lys	Thr	Phe	Gln	Val
				575					580					585
Leu	Gly	Asn	Leu	Tyr	Ser	Glu	Gly	Asp	Cys	Thr	Tyr	Leu	Lys	Cys
				590					595					600
Ser	Val	Asn	Gly	Val	Ala	Ser	Lys	Ala	Lys	Leu	Ile	Ile	Leu	Glu

Asn Thr Ile Tyr	605	610	615
Leu Phe Ser Lys Glu Gly Ser Ile Glu Ile Asp	620	625	630
Ile Pro Val Pro Lys Tyr Leu Ser Ser Val Ser Ser Gln Glu Thr	635	640	645
Gln Gly Gly Pro Leu Ala Pro Met Thr Gly Thr Ile Glu Lys Val	650	655	660
Phe Val Lys Ala Gly Asp Lys Val Lys Ala Gly Asp Ser Leu Met	665	670	675
Val Met Ile Ala Met Lys Met Glu His Thr Ile Lys Ser Pro Lys	680	685	690
Asp Gly Thr Val Lys Lys Val Phe Tyr Arg Glu Gly Ala Gln Ala	695	700	705
Asn Arg His Thr Pro Leu Val Glu Phe Glu Glu Glu Glu Ser Asp	710	715	720
Lys Arg Glu Ser Glu	725		

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<211> 644

<212> PRT

<213> Homo sapiens

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<223> Incyte ID No: 2844928CD1

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Met Ala Val Tyr Val Gly Met Leu Arg Leu Gly Arg Leu Cys Ala	1	5	10	15
Gly Ser Ser Gly Val Leu Gly Ala Arg Ala Leu Ser Arg Ser	20	25	30	35
Trp Gln Glu Ala Arg Leu Gln Gly Val Arg Phe Leu Ser Ser Arg	40	45	50	55
Glu Val Asp Arg Met Val Ser Thr Pro Ile Gly Gly Leu Ser Tyr	60	65	70	75
Val Gln Gly Cys Thr Lys Lys His Leu Asn Ser Lys Thr Val Gly	80	85	90	95
Gln Cys Leu Glu Thr Thr Ala Gln Arg Val Pro Glu Arg Glu Ala	100	105	110	115
Leu Val Val Leu His Glu Asp Val Arg Leu Thr Phe Ala Gln Leu	120	125	130	135
Lys Glu Glu Val Asp Lys Ala Ala Ser Gly Leu Leu Ser Ile Gly	140	145	150	155
Leu Cys Lys Gly Asp Arg Leu Gly Met Trp Gly Pro Asn Ser Tyr	160	165	170	175
Ala Trp Val Leu Met Gln Leu Ala Thr Ala Gln Ala Gly Ile Ile	180	185	190	195
Leu Val Ser Val Asn Pro Ala Tyr Gln Ala Met Glu Leu Glu Tyr	200	205	210	215
Val Leu Lys Lys Val Gly Cys Lys Ala Leu Val Phe Pro Lys Gln	220	225	230	235
Phe Lys Thr Gln Gln Tyr Tyr Asn Val Leu Lys Gln Ile Cys Pro	240	245	250	255
Glu Val Glu Asn Ala Gln Pro Gly Ala Leu Lys Ser Gln Arg Leu	260	265	270	275
Pro Asp Leu Thr Thr Val Ile Ser Val Asp Ala Pro Leu Pro Gly	280			
Thr Leu Leu Leu Asp Glu Val Val Ala Ala Gly Ser Thr Arg Gln				
His Leu Asp Gln Leu Gln Tyr Asn Gln Gln Phe Leu Ser Cys His				
Asp Pro Ile Asn Ile Gln Phe Thr Ser Gly Thr Thr Gly Ser Pro				
Lys Gly Ala Thr Leu Ser His Tyr Asn Ile Val Asn Asn Ser Asn				

Ile	Leu	Gly	Glu	Arg	Leu	Lys	Leu	His	Glu	Lys	Thr	Pro	Glu	Gln	290	295	300
Leu	Arg	Met	Ile	Leu	Pro	Asn	Pro	Leu	Tyr	His	Cys	Leu	Gly	Ser	305	310	315
Val	Ala	Gly	Thr	Met	Met	Cys	Leu	Met	Tyr	Gly	Ala	Thr	Leu	Ile	320	325	330
Leu	Ala	Ser	Pro	Ile	Phe	Asn	Gly	Lys	Lys	Ala	Leu	Glu	Ala	Ile	335	340	345
Ser	Arg	Glu	Arg	Gly	Thr	Phe	Leu	Tyr	Gly	Thr	Pro	Thr	Met	Phe	350	355	360
Val	Asp	Ile	Leu	Asn	Gln	Pro	Asp	Phe	Ser	Ser	Tyr	Asp	Ile	Ser	365	370	375
Thr	Met	Cys	Gly	Gly	Val	Ile	Ala	Gly	Ser	Pro	Ala	Pro	Pro	Glu	380	385	390
Leu	Ile	Arg	Ala	Ile	Ile	Asn	Lys	Ile	Asn	Met	Lys	Asp	Leu	Val	395	400	405
Val	Ala	Tyr	Gly	Thr	Thr	Glu	Asn	Ser	Pro	Val	Thr	Phe	Ala	His	410	415	420
Phe	Pro	Glu	Asp	Thr	Val	Glu	Gln	Lys	Ala	Glu	Ser	Val	Gly	Arg	425	430	435
Ile	Met	Pro	His	Thr	Glu	Ala	Arg	Ile	Met	Asn	Met	Glu	Ala	Gly	440	445	450
Thr	Leu	Ala	Lys	Leu	Asn	Thr	Pro	Gly	Glu	Leu	Cys	Ile	Arg	Gly	455	460	465
Tyr	Cys	Val	Met	Leu	Gly	Tyr	Trp	Gly	Glu	Pro	Gln	Lys	Thr	Glu	470	475	480
Glu	Ala	Val	Asp	Gln	Asp	Lys	Trp	Tyr	Trp	Thr	Gly	Asp	Val	Ala	485	490	495
Thr	Met	Asn	Glu	Gln	Gly	Phe	Cys	Lys	Ile	Val	Gly	Arg	Ser	Lys	500	505	510
Asp	Met	Ile	Ile	Arg	Gly	Gly	Glu	Asn	Ile	Tyr	Pro	Ala	Glu	Leu	515	520	525
Glu	Asp	Phe	Phe	His	Thr	His	Pro	Lys	Val	Gln	Glu	Val	Gln	Val	530	535	540
Arg	His	Leu	Ala	Gln	Val	Ser	Pro	Gln	Lys	Gln	Glu	Thr	His	Met	545	550	555
Asn	Thr	Val	Met	Ser	Asp	Ile	Phe	Leu	Trp	Pro	Trp	Asn	Val	Val	560	565	570
Gly	Val	Lys	Asp	Asp	Arg	Met	Gly	Glu	Glu	Ile	Cys	Ala	Cys	Ile	575	580	585
Arg	Leu	Lys	Asp	Gly	Glu	Glu	Thr	Thr	Val	Glu	Glu	Ile	Lys	Ala	590	595	600
Phe	Cys	Lys	Gly	Lys	Ile	Ser	His	Phe	Lys	Ile	Pro	Lys	Tyr	Ile	605	610	615
Val	Phe	Val	Thr	Asn	Tyr	Pro	Leu	Thr	Ile	Ser	Gly	Lys	Ile	Gln	620	625	630
Lys	Phe	Lys	Leu	Arg	Glu	Gln	Met	Glu	Arg	His	Leu	Asn	Leu		635	640	

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<211> 504

<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 3231586CD1

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Gly	Phe	Leu	Gly	Val	Tyr	Tyr	Val	Gly	Val	Ala	Ser	Cys	Leu	Arg	20	25	30	
Glu	His	Ala	Pro	Phe	Leu	Val	Ala	Asn	Ala	Thr	His	Ile	Tyr	Gly	35	40	45	
Ala	Ser	Ala	Gly	Ala	Leu	Thr	Ala	Thr	Ala	Leu	Val	Thr	Gly	Val				

	50		55		60									
Cys	Leu	Gly	Glu	Ala	Gly	Ala	Lys	Phe	Ile	Glu	Val	Ser	Lys	Glu
	65		70		75									
Ala	Arg	Lys	Arg	Phe	Leu	Gly	Pro	Leu	His	Pro	Ser	Phe	Asn	Leu
	80		85		90									
Val	Lys	Ile	Ile	Arg	Ser	Phe	Leu	Leu	Lys	Val	Leu	Pro	Ala	Asp
	95		100		105									
Ser	His	Glu	His	Ala	Ser	Gly	Arg	Leu	Gly	Ile	Ser	Leu	Thr	Arg
	110		115		120									
Val	Ser	Asp	Gly	Glu	Asn	Val	Ile	Ile	Ser	His	Phe	Asn	Ser	Lys
	125		130		135									
Asp	Glu	Leu	Ile	Gln	Ala	Asn	Val	Cys	Ser	Gly	Phe	Ile	Pro	Val
	140		145		150									
Tyr	Cys	Gly	Leu	Ile	Pro	Pro	Ser	Leu	Gln	Gly	Val	Arg	Tyr	Val
	155		160		165									
Asp	Gly	Gly	Ile	Ser	Asp	Asn	Leu	Pro	Leu	Tyr	Glu	Leu	Lys	Asn
	170		175		180									
Thr	Ile	Thr	Val	Ser	Pro	Phe	Ser	Gly	Glu	Ser	Asp	Ile	Cys	Pro
	185		190		195									
Gln	Asp	Ser	Ser	Thr	Asn	Ile	His	Glu	Leu	Arg	Val	Thr	Asn	Thr
	200		205		210									
Ser	Ile	Gln	Phe	Asn	Leu	Arg	Asn	Leu	Tyr	Arg	Leu	Ser	Lys	Ala
	215		220		225									
Leu	Phe	Pro	Pro	Glu	Pro	Leu	Val	Leu	Arg	Glu	Met	Cys	Lys	Gln
	230		235		240									
Gly	Tyr	Arg	Asp	Gly	Leu	Arg	Phe	Leu	Gln	Arg	Asn	Gly	Leu	Leu
	245		250		255									
Asn	Arg	Pro	Asn	Pro	Leu	Leu	Ala	Leu	Pro	Pro	Ala	Arg	Pro	His
	260		265		270									
Gly	Pro	Glu	Asp	Lys	Asp	Gln	Ala	Val	Glu	Ser	Ala	Gln	Ala	Glu
	275		280		285									
Asp	Tyr	Ser	Gln	Leu	Pro	Gly	Glu	Asp	His	Ile	Leu	Glu	His	Leu
	290		295		300									
Pro	Ala	Arg	Leu	Asn	Glu	Ala	Leu	Leu	Glu	Ala	Cys	Val	Glu	Pro
	305		310		315									
Thr	Asp	Leu	Leu	Thr	Thr	Leu	Ser	Asn	Met	Leu	Pro	Val	Arg	Leu
	320		325		330									
Ala	Thr	Ala	Met	Met	Val	Pro	Tyr	Thr	Leu	Pro	Leu	Glu	Ser	Ala
	335		340		345									
Leu	Ser	Phe	Thr	Ile	Arg	Leu	Leu	Glu	Trp	Leu	Pro	Asp	Val	Pro
	350		355		360									
Glu	Asp	Ile	Arg	Trp	Met	Lys	Glu	Gln	Thr	Gly	Ser	Ile	Cys	Gln
	365		370		375									
Tyr	Leu	Val	Met	Arg	Ala	Lys	Arg	Lys	Leu	Gly	Arg	His	Leu	Pro
	380		385		390									
Ser	Arg	Leu	Pro	Glu	Gln	Val	Glu	Leu	Arg	Arg	Val	Gln	Ser	Leu
	395		400		405									
Pro	Ser	Val	Pro	Leu	Ser	Cys	Ala	Ala	Tyr	Arg	Glu	Ala	Leu	Pro
	410		415		420									
Gly	Trp	Met	Arg	Asn	Asn	Leu	Ser	Leu	Gly	Asp	Ala	Leu	Ala	Lys
	425		430		435									
Trp	Glu	Glu	Cys	Gln	Arg	Gln	Leu	Leu	Leu	Gly	Leu	Phe	Cys	Thr
	440		445		450									
Asn	Val	Ala	Phe	Pro	Pro	Glu	Ala	Leu	Arg	Met	Arg	Ala	Pro	Ala
	455		460		465									
Asp	Pro	Ala	Pro	Ala	Pro	Ala	Asp	Pro	Ala	Ser	Pro	Gln	His	Gln
	470		475		480									
Leu	Ala	Gly	Pro	Ala	Pro	Leu	Leu	Ser	Thr	Pro	Ala	Pro	Glu	Ala
	485		490		495									
Arg	Pro	Val	Ile	Gly	Ala	Leu	Gly	Leu						
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 <213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 3580770CD1

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Lys	Gly	Ile	Ile	Ala	Ser	Ser	Ile	Gly	Thr	Ile	Leu	Lys	Ser	Cys
				20					25					30
Gly	Leu	Arg	Val	Thr	Ala	Ile	Lys	Ile	Asp	Pro	Tyr	Ile	Asn	Ile
				35					40					45
Asp	Ala	Gly	Thr	Phe	Ser	Pro	Tyr	Glu	His	Gly	Glu	Val	Phe	Val
				50					55					60
Leu	Asn	Asp	Gly	Gly	Glu	Val	Asp	Leu	Asp	Leu	Gly	Asp	Tyr	Glu
				65					70					75
Arg	Phe	Leu	Asp	Ile	Asn	Leu	Tyr	Lys	Asp	Thr	Ile	Val	Thr	Thr
				80					85					90
Gly	Lys	Ile	Tyr	Gln	His	Val	Ile	Asn	Lys	Glu	Arg	Arg	Gly	Asp
				95					100					105
Tyr	Leu	Gly	Lys	Thr	Val	Gln	Val	Val	Pro	His	Ile	Thr	Asp	Ala
				110					115					120
Val	Gln	Glu	Trp	Val	Met	Asn	Gln	Ala	Lys	Val	Pro	Val	Asp	Gly
				125					130					135
Asn	Lys	Glu	Glu	Pro	Gln	Ile	Cys	Val	Ile	Glu	Leu	Gly	Gly	Thr
				140					145					150
Ile	Gly	Asp	Ile	Glu	Gly	Met	Pro	Phe	Val	Glu	Ala	Phe	Arg	Gln
				155					160					165
Phe	Gln	Phe	Lys	Ala	Lys	Arg	Glu	Asn	Phe	Cys	Asn	Ile	His	Val
				170					175					180
Ser	Leu	Val	Pro	Gln	Leu	Ser	Ala	Thr	Gly	Glu	Gln	Lys	Thr	Lys
				185					190					195
Pro	Thr	Gln	Asn	Ser	Val	Arg	Ala	Leu	Arg	Gly	Leu	Gly	Leu	Ser
				200					205					210
Pro	Asp	Leu	Ile	Val	Cys	Arg	Ser	Ser	Thr	Pro	Ile	Glu	Met	Ala
				215					220					225
Val	Lys	Glu	Lys	Ile	Ser	Met	Phe	Cys	His	Val	Asn	Pro	Glu	Gln
				230					235					240
Val	Ile	Cys	Ile	His	Asp	Val	Ser	Ser	Thr	Tyr	Arg	Val	Pro	Val
				245					250					255
Leu	Leu	Glu	Glu	Gln	Ser	Ile	Val	Lys	Tyr	Phe	Lys	Glu	Arg	Leu
				260					265					270
His	Leu	Pro	Ile	Gly	Asp	Ser	Ala	Ser	Asn	Leu	Leu	Phe	Lys	Trp
				275					280					285
Arg	Asn	Met	Ala	Asp	Arg	Tyr	Glu	Arg	Leu	Gln	Lys	Ile	Cys	Ser
				290					295					300
Ile	Ala	Leu	Val	Gly	Lys	Tyr	Thr	Lys	Leu	Arg	Asp	Cys	Tyr	Ala
				305					310					315
Ser	Val	Phe	Lys	Ala	Leu	Glu	His	Ser	Ala	Leu	Ala	Ile	Asn	His
				320					325					330
Lys	Leu	Asn	Leu	Met	Val	Ile	Asp	Met	Pro	Glu	His	Asn	Pro	Gly
				335					340					345
Asn	Leu	Gly	Gly	Thr	Met	Arg	Leu	Gly	Ile	Arg	Arg	Thr	Val	Phe
				350					355					360
Lys	Thr	Glu	Asn	Ser	Ile	Leu	Arg	Lys	Leu	Tyr	Gly	Asp	Val	Pro
				365					370					375
Phe	Ile	Glu	Glu	Arg	His	Arg	His	Arg	Phe	Glu	Val	Asn	Pro	Asn
				380					385					390
Leu	Ile	Lys	Gln	Phe	Glu	Gln	Asn	Asp	Leu	Ser	Phe	Val	Gly	Gln
				395					400					405
Asp	Val	Asp	Gly	Asp	Arg	Met	Glu	Ile	Ile	Glu	Leu	Ala	Asn	His
				410					415					420
Pro	Tyr	Phe	Val	Gly	Val	Gln	Phe	His	Pro	Glu	Phe	Ser	Ser	Arg
				425					430					435
Pro	Met	Lys	Pro	Ser	Pro	Pro	Tyr	Leu	Gly	Leu	Leu	Leu	Ala	Ala
				440					445					450

Thr Gly Asn Leu Asn Ala Tyr Leu Gln Gln Gly Cys Lys Leu Ser
 455 460 465
 Ser Ser Asp Arg Tyr Ser Asp Ala Ser Asp Asp Ser Phe Ser Glu
 470 475 480
 Pro Arg Ile Ala Glu Leu Glu Ile Ser
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 <211> 258
 <212> PRT
 <213> Homo sapiens

<220>
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 <223> Incyte ID No: 3778612CD1

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 Phe Ile Gln Ser Thr Leu Pro Leu Lys Gln Glu Glu Tyr Glu Ala
 20 25 30
 Phe Leu Leu Lys Leu Val Gln Asn Leu Phe Ala Glu Gly Asn Asp
 35 40 45
 Leu Phe Arg Glu Lys Asp Tyr Lys Gln Ala Leu Val Gln Tyr Met
 50 55 60
 Glu Gly Leu Asn Val Ala Asp Tyr Ala Ala Ser Asp Gln Val Ala
 65 70 75
 Leu Pro Arg Glu Leu Leu Cys Lys Leu His Val Asn Arg Ala Ala
 80 85 90
 Cys Tyr Phe Thr Met Gly Leu Tyr Glu Lys Ala Leu Glu Asp Ser
 95 100 105
 Glu Lys Ala Leu Gly Pro Asp Ser Glu Ser Ile Arg Ala Leu Phe
 110 115 120
 Arg Lys Ala Arg Ala Leu Asn Glu Leu Gly Arg His Lys Glu Ala
 125 130 135
 Tyr Glu Cys Ser Ser Arg Cys Ser Leu Ala Leu Pro His Asp Glu
 140 145 150
 Ser Val Thr Gln Leu Gly Gln Gly Pro Leu Gly Ser Gly Ala Ser
 155 160 165
 Trp Pro Gly Gln Ser Trp Ser Pro His Arg Val Arg Lys Arg Glu
 170 175 180
 Trp Glu Ala Glu Cys Asp Gly Glu Glu Gly Gln Glu Asp Pro Phe
 185 190 195
 Asn Asp Glu Gly Asn Tyr Phe Ser Cys Glu Pro Ser Arg Ala Pro
 200 205 210
 Gly Trp Glu Ala Gln Arg Thr Glu Ser Gly Thr Cys Val Pro Pro
 215 220 225
 Gly Arg Gln Gly Gln Asp Gly Met Ala Ser Met Gly Ala Gly Trp
 230 235 240
 Val Gly Arg Asp Ala Ala Phe Leu Ser Trp Ala Val Ile Asn Leu
 245 250 255
 Met Val Leu

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<220>
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 <222> 63-65
 <223> unknown or other

<400> 12

Met	Glu	Gly	Ala	Val	Leu	Glu	Ala	Gly	Gly	Ala	Arg	Cys	Phe	Cys	
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Arg	Phe	Gly	Cys	Glu	Leu	Ser	Lys	Tyr	Glu	Asn	Pro	Gly	Tyr	Ser	
				20					25					30	
Ser	Pro	Arg	Ser	Asp	Tyr	Phe	Lys	Asn	Tyr	Met	Ile	Ile	Ile	Thr	
				35					40					45	
Gln	Asn	Arg	Met	Ser	Phe	Leu	Ala	Asn	Met	Phe	His	Thr	Met	Asp	
				50					55					60	
Cys	Val	Xaa	Xaa	Xaa	Arg	Tyr	Ser	Cys	Gly	Pro	Thr	Val	Tyr	Asp	
				65					70					75	
His	Ala	His	Leu	Gly	His	Ala	Cys	Ser	Tyr	Val	Arg	Phe	Asp	Ile	
				80					85					90	
Ile	Arg	Arg	Ile	Leu	Thr	Lys	Val	Phe	Gly	Cys	Ser	Ile	Val	Met	
				95					100					105	
Val	Met	Gly	Ile	Thr	Asp	Val	Asp	Asp	Lys	Ile	Ile	Lys	Arg	Ala	
				110					115					120	
Asn	Glu	Met	Asn	Ile	Ser	Pro	Ala	Ser	Leu	Ala	Ser	Leu	Tyr	Glu	
				125					130					135	
Glu	Asp	Phe	Lys	Gln	Asp	Met	Ala	Ala	Leu	Lys	Val	Leu	Pro	Pro	
				140					145					150	
Thr	Val	Tyr	Leu	Arg	Val	Thr	Glu	Asn	Ile	Pro	Gln	Ile	Ile	Ser	
				155					160					165	
Phe	Ile	Glu	Gly	Ile	Ile	Ala	Arg	Gly	Asn	Ala	Tyr	Ser	Thr	Ala	
				170					175					180	
Lys	Gly	Asn	Val	Tyr	Phe	Asp	Leu	Lys	Ser	Arg	Gly	Asp	Lys	Tyr	
				185					190					195	
Gly	Lys	Leu	Val	Gly	Val	Val	Pro	Gly	Pro	Val	Gly	Glu	Pro	Ala	
				200					205					210	
Asp	Ser	Asp	Lys	Arg	His	Ala	Ser	Asp	Phe	Ala	Leu	Trp	Lys	Ala	
				215					220					225	
Ala	Lys	Pro	Gln	Glu	Val	Phe	Trp	Ala	Ser	Pro	Trp	Gly	Pro	Gly	
				230					235					240	
Arg	Pro	Gly	Trp	His	Ile	Glu	Cys	Ser	Ala	Ile	Ala	Ser	Met	Val	
				245					250					255	
Phe	Gly	Ser	Gln	Leu	Asp	Ile	His	Ser	Gly	Gly	Ile	Asp	Leu	Ala	
				260					265					270	
Phe	Pro	His	His	Glu	Asn	Glu	Ile	Ala	Gln	Cys	Glu	Val	Phe	His	
				275					280					285	
Gln	Cys	Glu	Gln	Trp	Gly	Asn	Tyr	Phe	Leu	His	Ser	Gly	His	Leu	
				290					295					300	
His	Ala	Lys	Gly	Lys	Glu	Glu	Lys	Met	Ser	Lys	Ser	Leu	Lys	Asn	
				305					310					315	
Tyr	Ile	Thr	Ile	Lys	Asp	Phe	Leu	Lys	Thr	Phe	Ser	Pro	Asp	Val	
				320					325					330	
Phe	Arg	Phe	Phe	Cys	Leu	Arg	Ser	Ser	Tyr	Arg	Ser	Ala	Ile	Asp	
				335					340					345	
Tyr	Ser	Asp	Ser	Ala	Met	Leu	Gln	Ala	Gln	Gln	Leu	Leu	Leu	Gly	
				350					355					360	
Leu	Gly	Ser	Phe	Leu	Glu	Asp	Ala	Arg	Ala	Tyr	Met	Lys	Gly	Gln	
				365					370					375	
Leu	Ala	Cys	Gly	Ser	Val	Arg	Glu	Ala	Met	Leu	Trp	Glu	Arg	Leu	
				380					385					390	
Ser	Ser	Thr	Lys	Arg	Ala	Val	Lys	Ala	Ala	Leu	Ala	Asp	Asp	Phe	
				395					400					405	
Asp	Thr	Pro	Arg	Val	Val	Asp	Ala	Ile	Leu	Gly	Leu	Ala	His	His	
				410					415					420	
Gly	Asn	Gly	Gln	Leu	Arg	Ala	Ser	Leu	Lys	Glu	Pro	Glu	Gly	Pro	
				425					430					435	
Arg	Ser	Pro	Ala	Val	Phe	Gly	Ala	Ile	Ile	Ser	Tyr	Phe	Glu	Gln	
				440					445					450	
Phe	Phe	Glu	Thr	Val	Gly	Ile	Ser	Leu	Ala	Asn	Gln	Gln	Tyr	Val	
				455					460					465	
Ser	Gly	Asp	Gly	Ser	Glu	Ala	Thr	Leu	His	Gly	Val	Val	Asp	Glu	
				470					475					480	

Leu	Val	Arg	Phe	Arg	Gln	Lys	Val	Arg	Gln	Phe	Ala	Leu	Ala	Met
				485					490					495
Pro	Glu	Ala	Thr	Gly	Asp	Ala	Arg	Arg	Gln	Gln	Leu	Leu	Glu	Arg
				500					505					510
Gln	Pro	Leu	Leu	Glu	Ala	Cys	Asp	Thr	Leu	Arg	Arg	Gly	Leu	Thr
				515					520					525
Ala	His	Gly	Ile	Asn	Ile	Lys	Asp	Arg	Ser	Ser	Thr	Thr	Ser	Thr
				530					535					540
Trp	Glu	Leu	Leu	Asp	Gln	Arg	Thr	Lys	Asp	Gln	Lys	Ser	Ala	Gly
				545					550					555

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<211> 463

<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 5630806CD1

<400> 13

Met	Ala	Ala	Ser	Met	Phe	Tyr	Gly	Arg	Leu	Val	Ala	Val	Ala	Thr
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Leu	Arg	Asn	His	Arg	Pro	Arg	Thr	Ala	Gln	Arg	Ala	Ala	Ala	Gln
				20					25					30
Val	Leu	Gly	Ser	Ser	Gly	Leu	Phe	Asn	Asn	His	Gly	Leu	Gln	Val
				35					40					45
Gln	Gln	Gln	Gln	Gln	Arg	Asn	Leu	Ser	Leu	His	Glu	Tyr	Met	Ser
				50					55					60
Met	Glu	Leu	Leu	Gln	Glu	Ala	Gly	Val	Ser	Val	Pro	Lys	Gly	Tyr
				65					70					75
Val	Ala	Lys	Ser	Pro	Asp	Glu	Ala	Tyr	Ala	Ile	Ala	Lys	Lys	Leu
				80					85					90
Gly	Ser	Lys	Asp	Val	Val	Ile	Lys	Ala	Gln	Val	Leu	Ala	Gly	Gly
				95					100					105
Arg	Gly	Lys	Gly	Thr	Phe	Glu	Ser	Gly	Leu	Lys	Gly	Gly	Val	Lys
				110					115					120
Ile	Val	Phe	Ser	Pro	Glu	Glu	Ala	Lys	Ala	Val	Ser	Ser	Gln	Met
				125					130					135
Ile	Gly	Lys	Lys	Leu	Phe	Thr	Lys	Gln	Thr	Gly	Glu	Lys	Gly	Arg
				140					145					150
Ile	Cys	Asn	Gln	Val	Leu	Val	Cys	Glu	Arg	Lys	Tyr	Pro	Arg	Arg
				155					160					165
Glu	Tyr	Tyr	Phe	Ala	Ile	Thr	Met	Glu	Arg	Ser	Phe	Gln	Gly	Pro
				170					175					180
Val	Leu	Ile	Gly	Ser	Ser	His	Gly	Gly	Val	Asn	Ile	Glu	Asp	Val
				185					190					195
Ala	Ala	Glu	Thr	Pro	Glu	Ala	Ile	Ile	Lys	Glu	Pro	Ile	Asp	Ile
				200					205					210
Glu	Glu	Gly	Ile	Lys	Lys	Glu	Gln	Ala	Leu	Gln	Leu	Ala	Gln	Lys
				215					220					225
Met	Gly	Phe	Pro	Pro	Asn	Ile	Val	Glu	Ser	Ala	Ala	Glu	Asn	Met
				230					235					240
Val	Lys	Leu	Tyr	Ser	Leu	Phe	Leu	Lys	Tyr	Asp	Ala	Thr	Met	Ile
				245					250					255
Glu	Ile	Asn	Pro	Met	Val	Glu	Asp	Ser	Asp	Gly	Ala	Val	Leu	Cys
				260					265					270
Met	Asp	Ala	Lys	Ile	Asn	Phe	Asp	Ser	Asn	Ser	Ala	Tyr	Arg	Gln
				275					280					285
Lys	Lys	Ile	Phe	Asp	Leu	Gln	Asp	Trp	Thr	Gln	Glu	Asp	Glu	Arg
				290					295					300
Asp	Lys	Asp	Ala	Ala	Lys	Ala	Asn	Leu	Asn	Tyr	Ile	Gly	Leu	Asp
				305					310					315
Gly	Asn	Ile	Gly	Cys	Leu	Val	Asn	Gly	Ala	Gly	Leu	Ala	Met	Ala
				320					325					330

Thr	Met	Asp	Ile	Ile	Lys	Leu	His	Gly	Gly	Thr	Pro	Ala	Asn	Phe
				335					340					345
Leu	Asp	Val	Gly	Gly	Gly	Ala	Thr	Val	His	Gln	Val	Thr	Glu	Ala
				350					355					360
Phe	Lys	Leu	Ile	Thr	Ser	Asp	Lys	Lys	Val	Leu	Ala	Ile	Leu	Val
				365					370					375
Asn	Ile	Phe	Gly	Gly	Ile	Met	Arg	Cys	Asp	Val	Ile	Ala	Gln	Gly
				380					385					390
Ile	Val	Met	Ala	Val	Lys	Asp	Leu	Glu	Ile	Lys	Ile	Pro	Val	Val
				395					400					405
Val	Arg	Leu	Gln	Gly	Thr	Arg	Val	Asp	Asp	Ala	Lys	Ala	Leu	Ile
				410					415					420
Ala	Asp	Ser	Gly	Leu	Lys	Ile	Leu	Ala	Cys	Asp	Asp	Leu	Asp	Glu
				425					430					435
Ala	Ala	Arg	Met	Val	Val	Lys	Leu	Ser	Glu	Ile	Val	Thr	Leu	Ala
				440					445					450
Lys	Gln	Ala	His	Val	Asp	Val	Lys	Phe	Gln	Leu	Pro	Ile		
				455					460					

<210> 14

<211> 399

<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 5854855CD1

<400> 14

Met	Tyr	Phe	Gly	Ala	Val	Ser	Trp	Leu	Cys	Gly	Pro	Arg	Ala	Pro
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Asp	Glu	Val	Ser	Arg	Arg	Pro	Asp	Pro	Arg	Lys	Gly	Gln	Leu	Gly
				20					25					30
Val	Ala	Phe	Val	Leu	Leu	Pro	Pro	His	Ser	Glu	Gly	Ala	Arg	Val
				35					40					45
Phe	Gly	Ala	Leu	Gly	Pro	Ile	Gly	Pro	Ser	Ser	Pro	Gly	Leu	Thr
				50					55					60
Leu	Gly	Gly	Leu	Ala	Val	Ser	Glu	His	Arg	Leu	Ser	Asn	Lys	Leu
				65					70					75
Leu	Ala	Trp	Ser	Gly	Val	Leu	Glu	Trp	Gln	Glu	Lys	Arg	Arg	Pro
				80					85					90
Tyr	Ser	Asp	Ser	Thr	Ala	Lys	Leu	Lys	Arg	Thr	Leu	Pro	Cys	Gln
				95					100					105
Ala	Tyr	Val	Asn	Gln	Gly	Glu	Asn	Leu	Glu	Thr	Asp	Gln	Trp	Pro
				110					115					120
Gln	Lys	Leu	Ile	Met	Gln	Leu	Ile	Pro	Gln	Gln	Leu	Leu	Thr	Thr
				125					130					135
Leu	Gly	Pro	Leu	Phe	Arg	Asn	Ser	Gln	Leu	Ala	Gln	Phe	His	Phe
				140					145					150
Thr	Asn	Arg	Asp	Cys	Asp	Ser	Leu	Lys	Gly	Leu	Cys	Arg	Ile	Met
				155					160					165
Gly	Asn	Gly	Phe	Ala	Gly	Cys	Met	Leu	Phe	Pro	His	Ile	Ser	Pro
				170					175					180
Cys	Glu	Val	Arg	Val	Leu	Met	Leu	Leu	Tyr	Ser	Ser	Lys	Lys	Lys
				185					190					195
Ile	Phe	Met	Gly	Leu	Ile	Pro	Tyr	Asp	Gln	Ser	Gly	Phe	Val	Ser
				200					205					210
Ala	Ile	Arg	Gln	Val	Ile	Thr	Thr	Arg	Lys	Gln	Ala	Val	Gly	Pro
				215					220					225
Gly	Gly	Val	Asn	Ser	Gly	Pro	Val	Gln	Ile	Val	Asn	Asn	Lys	Phe
				230					235					240
Leu	Ala	Trp	Ser	Gly	Val	Met	Glu	Trp	Gln	Glu	Pro	Arg	Pro	Glu
				245					250					255
Pro	Asn	Ser	Arg	Ser	Lys	Arg	Trp	Leu	Pro	Ser	His	Val	Tyr	Val
				260					265					270
Asn	Gln	Gly	Glu	Ile	Leu	Arg	Thr	Glu	Gln	Trp	Pro	Arg	Lys	Leu

	275		280		285
Tyr Met Gln Leu	Ile Pro Gln Gln Leu	Leu Thr Thr Leu Val	Pro		
	290		295		300
Leu Phe Arg Asn	Ser Arg Leu Val Gln	Phe His Phe Thr Lys	Asp		
	305		310		315
Leu Glu Thr Leu	Lys Ser Leu Cys Arg	Ile Met Asp Asn Gly	Phe		
	320		325		330
Ala Gly Cys Val	His Phe Ser Tyr Lys	Ala Ser Cys Glu Ile	Arg		
	335		340		345
Val Leu Met Leu	Leu Tyr Ser Ser Glu	Lys Lys Ile Phe Ile	Gly		
	350		355		360
Leu Ile Pro His	Asp Gln Gly Asn Phe	Val Asn Gly Ile Arg	Arg		
	365		370		375
Val Ile Ala Asn	Gln Gln Gln Val Leu	Gln Arg Asn Leu Glu	Gln		
	380		385		390
Glu Gln Gln Gln	Arg Gly Met Gly Gly				
	395				

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<211> 339

<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 5993973CD1

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Val Gly Gly Arg Gly	Gly Gly Lys Gly	Lys Gly Arg Ala Glu	Gly		
	20	25	30		
Ser Glu Lys Glu Glu	Ser Arg Arg Lys	Arg Arg Glu Arg Lys	Gln		
	35	40	45		
Arg Arg Glu Gly Gly	Asp Gly Glu Glu	Gln Asp Val Gly Asp	Ala		
	50	55	60		
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<213> Homo sapiens

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<220>

<221> misc_feature

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<212> DNA
<213> Homo sapiens

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<211> 1954
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<213> Homo sapiens

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<211> 1937

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<210> 26

<211> 970

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 3778612CB1

<400> 26

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<210> 27

<211> 1810

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 4574912CB1

<220>

<221> unsure

<222> 193, 196-198

<223> a, t, c, g, or other

<400> 27

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<211> 2162
<212> DNA
<213> Homo sapiens

<220>
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aa 2162

<210> 29
<211> 1477
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 5854855CB1

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<211> 1660
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
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